SBW2019 Schedule

	Monday 21 Oct.	
9:00 - 10:00	Registration and welcome + coffee (Set up posters)	
10:00 - 10:05	Opening address	
10:05 - 11:00	Christina Leslie Decoding epigenomic programs in cancer and immunity	
11:00 - 11:30	Nikolay Oskolkov National Bioinformatics Infrastructure Sweden (NBIS)	
11:30 - 12:00	Christoph Börlin Modeling TF binding profiles through competitive binding	
12:00 - 12:30	Fredrik Karlsson Bioinformatics and machine learning for generation and characterization of advanced cell models for drug discovery	
12:30 - 13:30	Lunch	
13:30 - 14:30	Lior Pachter Some solved and unsolved bioinformatics problems in single- cell genomics	
14:30 - 15:00	Aleksej Zelezniak Uncovering DNA grammar using deep learning	
15:00 - 16:00	Coffee break + Poster session 1	
16:00 - 18:00	Anna Reymer Workshop Understanding the biological function of a protein using structural bioinformatics	Johan Bengtsson-Palme Workshop Wait a minute! – On annotation errors and how to spot them
18:00	Dinner (at Wallenberg Conference Center)	

	Tuesday 22 Oct.	
9:00 - 10:00	Mihaela Zavolan Learning miRNA-dependent regulatory networks from high- throughput data	
10:00 - 10:30	Sara Younes Identification of combinatorial markers in Systemic Lupus Erythematosus and disease subtypes	
10:30 - 11:00	Coffee break + Poster session 2	
11:00 - 11:30	Olukayode Daramola Differential Expression of Annotated Adenylate Cyclase Genes in various Life Stages of Fasciola hepatica	
11:30 - 12:00	Marcela Davila University of Gothenburg Bioinformatics Core Facility	
12:00 - 12:30	Johanna Hörberg Regulatory role of torsional stress on bZIP transcription factors Interactions with DNA	
12:30 - 13:30	Lunch	
13:30 - 14:00	Francesco Gatto Elypta	
14:00 - 14:30	Erik Larsson Lekholm Understanding and modeling regional mutational heterogeneity in skin cancers	
14:30 - 15:00	Coffee break	
15:00 - 16:00	Careers in Computational Biology (hosted by <u>RSG-Sweden</u>) [http://www.rsg-sweden.iscbsc.org/]	
16:00 - 16:05	Closing remarks	

Legend

